## Claims

- A recombinant DNA sequence encoding an analog of mammalian FGF.
- 2. The DNA sequence of claim 1 which encodes a human FGF protein analog.
- 3. The DNA sequence of claim 2 which encodes a 10 human basic FGF protein analog.
  - 4. The DNA sequence of claim 3 which encodes a human basic FGF protein analog with reduced affinity for heparin binding.
- 5. The DNA sequence of claim 4 encoding a human basic FGF protein analog, comprising substituting one or more positively charged amino acid residues located in a heparin binding domain encompassing residues 128 through 138 with a neutral or negatively charged amino acid.
  - 6. The DNA sequence of claim 5 wherein the neutral or negatively charged amino acid is selected from the group consisting of serine, threonine or glutamic acid.
  - 7. The DNA sequence of claim 5 wherein the location and composition of the substituted amino acid is selected from the group consisting of serine 128, glutamic acid 128, threonine 129, serine 128/threonine 129, and serine 138.

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- 8. The DNA sequence of claim 2 which encodes a human basic FGF protein analog wherein one or more cysteine residues are replaced by a neutral amino acid and said protein analog exhibits the biological activity of native basic FGF.
- 9. The DNA sequence of claim 8 wherein the neutral amino acid is serine or alamine.
- 10. The DNA sequence of claim 9 wherein the substituted cysteine residue is at position 78, 96 or a combination thereof.
- 11. The DNA sequence of claim 5 which encodes a human basic FGF protein analog wherein said analog binds to a receptor for FGF and has reduced ability to induce a biological response.
- 12. The DNA sequence of claim 3 which encodes 20 an amino-terminal deletion analog of FGF having FGF antagonist activity.
  - 13. The DNA sequence of claim 12 wherein said deletion spans residues 1 through 24 of human basic FGF.
  - 14. The DNA sequence of claim 12 encoding a human basic FGF analog further comprising one or more positively charged amino acid residues located in a heparin binding domain encompassing residues 128 through 138 substituted with a neutral or negatively charged amino acid.
  - 15. The DNA sequence of claim 3 which is operably linked to control sequences for expression.

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- 16. The DNA sequence of claim 15 wherein the control sequences include a transcription termination signal.
- 5 17. The DNA sequence of claim 3 which is transformed into a recombinant host cell.
- 18. A recombinant vector containing the DNA sequence of claim 3 and effective in expressing FGF or an analog thereof.
  - 19. The vector of claim 18 which is selected from the group consisting of plasmids pUC9-TSF11 and pUC9delH3-pTSF-3.
  - 20. The vector of claim 18 wherein the DNA sequence encoding an FGF analog is operably linked to control sequences compatible with bacteria.
- 21. The vector of claim 18 wherein the DNA sequence encoding an FGF analog is operably linked to control sequences compatible with mammalian hosts.
- 22. Recombinant host cells transformed with the vector of claim 18.
  - $\,$  23. Bacterial cells transformed with the vector of claim 20.
- 30 24. Mammalian cells transformed with the vector of claim 21.
- 25. A method for producing FGF protein analogs which comprises culturing host cells harboring the DNA of claim 3 and recovering the FGF protein analog.

- 27. The method of claim 25 wherein the host 5 cells are mammalian.
- 28. A human basic FGF protein analog having reduced affinity for heparin binding comprising substituting one or more positively charged amino acid residues located in a heparin binding domain encompassing residues 128 through 138 with a neutral or negatively charged amino acid.
- 29. A human basic FGF protein analog wherein the cysteine at positions 78, 96, or a combination thereof, is replaced by a neutral amino acid and said analog exhibits the biological activity of native, human basic FGF.
- 20 30. The human basic FGF protein analog of claim 29 which is bFGF-C78/96S.
  - 31. An antagonist of human basic FGF.
- 32. The FCF antagonist of claim 31 wherein the first 24 amino terminal residues of basic FGF are deleted.

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